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**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE**

In re patent application of:

Berens, *et al.*

Appl. No.: 09/852,053

Filed: May 10, 2001

For: **Corynebacterium glutamicum  
strain with enhanced secretion  
activity**

Art Unit: to be assigned

Examiner: to be assigned

Atty. Dkt. 21123/280248

**Amendment to Comply With Sequence Listing Rules**

Assistant Commissioner for Patents  
Washington, D.C. 20231

Sir:

In response to the Notice to File Missing Parts dated June 4, 2001, Applicants are submitting the following amendments and remarks to comply with the Sequence Listing rules.

**Amendments**

Please delete the Sequence Listing originally filed with the application and appearing as pages 25-30 of the specification. Please enter the substitute Sequence Listing enclosed herewith on separate pages following the claims and abstract of the application.

On page 17 of the specification, please amend the table that starts on line 21 to read as follows:

| Gene          | 5'-primer   | 3'-primer   |
|---------------|---|---|
| secA fragment | saf1: 5'-CGCGACAAGGACTACATCGT-3'<br>(SEQ ID NO: 5)  | saf2: 5'-GAGATGTCTGCGGATTCGAG-3'<br>(SEQ ID NO: 15)   |
| secY fragment | syf1: 5'-TGAGGAGGCCAGGAGGCCAG-3'<br>(SEQ ID NO: 6)  | syf2: 5'-AACCAACCGTACTGACGACGA-3'<br>(SEQ ID NO: 16)  |
| secY          | sy1: 5'-TTAAGTGCTGAGGAGGCCAG-3'<br>(SEQ ID NO: 7)   | sy2: 5'-TTATCAGCACCGTAGTTCC-3'<br>(SEQ ID NO: 17)     |
| secE          | se1: 5'-TGGATGAGTAGTGATTAGA-3'<br>(SEQ ID NO: 8)    | se2: 5'-GATTCTGACTCCGTAGGTAG-3'<br>(SEQ ID NO: 18)    |
| secE region   | ser1: 5'-CACCTGGCAGACGCCTCAA-3'<br>(SEQ ID NO: 9)   | ser2: 5'-AGCCGGAGTAGCACTGAATG-3'<br>(SEQ ID NO: 19)   |
| secG          | sg1: 5'- ACCTGGGTTCTCAAACGGCA-3'<br>(SEQ ID NO: 10) | sg2: 5'-TTGTCGACCTGTTGTCTCCC-3'<br>(SEQ ID NO: 20)    |
| secG region   | sgr1: 5'-TCCAGGCCTTCCTCACGCAA-3'<br>(SEQ ID NO: 11) | sgr2: 5'-AGCTGCGAGAACATCCAGGCTA-3'<br>(SEQ ID NO: 21) |
| secD          | sd1: 5'-TTGTCTGGTTGATTGGAATT-3'<br>(SEQ ID NO: 12)  | sd2: 5'-TGAAGTTTCAGTCTGGGAAT-3'<br>(SEQ ID NO: 22)    |
| secD fragment | sdf1: 5'-TGCTGTTGACAGGCGATCGT-3'<br>(SEQ ID NO: 13) | sdf2: 5'-TCATCAGTGGTGCCTGCAT-3'<br>(SEQ ID NO: 23)    |
| secF fragment | sff1: 5'-GTACCAAGATGAGCATGCCA-3'<br>(SEQ ID NO: 14) | sff2: 5'-ATCGAACGCATGAAGGTCTG-3'<br>(SEQ ID NO: 24)   |